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SEQUENCE LISTING

<110> SUGIMOTO, Yukiko

ASAKURA, Yoko

MORI, Yukiko

ITO, Hisao

KURAHASHI, Osamu

<120> ARGININE REPRESSOR DEFICIENT STRAIN OF CORYNEFORM BACTERIUM AND METHOD FOR PRODUCING L-ARGININE

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Ala Val Ser Trp Ser Ser Asn Gly Ala Ser Gln Phe Glu Gly Leu Gln

50 55 60

Asn Ser His Ser Arg Trp Gly Ser Ser Leu Ala Glu Leu Glu Val Met
65 70 75 80

Gly Glu Arg Arg Ile Glu Leu Ala Ile Ala Thr Lys Asn His Leu Ala
85 90 95

Ala Gly Gly Ala Leu Met Met Phe Val Gly Thr Val Arg His Asn Arg
100 105 110

Ser Gln Ser Phe Ala Gln Val Glu Ala Gly Ile Lys Thr Ala Tyr Ser
115 120 125

Ser Met Val Lys Thr Ser Gln Trp Lys Lys Glu Arg Ala Arg Tyr Gly
130 135 140

Val Glu His Thr Tyr Ser Asp Tyr Glu Val Thr Asp Ser Trp Ala Asn
145 150 155 160

Gly Trp His Leu His Arg Asn Met Leu Leu Phe Leu Asp Arg Pro Leu
165 170 175

Ser Asp Asp Glu Leu Lys Ala Phe Glu Asp Ser Met Phe Ser Arg Trp
180 185 190

Ser Ala Gly Val Val Lys Ala Gly Met Asp Ala Pro Leu Arg Glu His
195 200 205

Gly Val Lys Leu Asp Gln Val Ser Thr Trp Gly Gly Asp Ala Ala Lys
210 215 220

Met Ala Thr Tyr Leu Ala Lys Gly Met Ser Gln Glu Leu Thr Gly Ser
225 230 235 240

Ala Thr Lys Thr Ala Ser Lys Gly Ser Tyr Thr Pro Phe Gln Met Leu
245 250 255

Asp Met Leu Ala Asp Gln Ser Asp Ala Gly Glu Asp Met Asp Ala Val

260 265 270

Leu Val Ala Arg Trp Arg Glu Tyr Glu Val Gly Ser Lys Asn Leu Arg
275 280 285

Ser Ser Trp Ser Arg Gly Ala Lys Arg Ala Leu Gly Ile Asp Tyr Ile
290 295 300

Asp Ala Asp Val Arg Arg Glu Met Glu Glu Glu Leu Tyr Lys Leu Ala
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Gly Leu Glu Ala Pro Glu Arg Val Glu Ser Thr Arg Val Ala Val Ala
325 330 335

Leu Val Lys Pro Asp Asp Trp Lys Leu Ile Gln Ser Asp Phe Ala Val
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Arg Gln Tyr Val Leu Asp Cys Val Asp Lys Ala Lys Asp Val Ala Ala
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Ala Gln Arg Val Ala Asn Glu Val Leu Ala Ser Leu Gly Val Asp Ser
370 375 380

Thr Pro Cys Met Ile Val Met Asp Asp Val Asp Leu Asp Ala Val Leu
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gcggtgtgga gttctaccct tacggcgaca ccgattactt gcgcaaaatg gtagaaacca 240

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                                                          Met Ser
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ctt ggc tca acc ccg tca aca ccg gaa aac tta aat ccc gtg act cgc      1905
Leu Gly Ser Thr Pro Ser Thr Pro Glu Asn Leu Asn Pro Val Thr Arg
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act gca cgc caa gct ctc att ttg cag att ttg gac aaa caa aaa gtc      1953
Thr Ala Arg Gln Ala Leu Ile Leu Gln Ile Leu Asp Lys Gln Lys Val
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acc agc cag gta caa ctg tct gaa ttg ctg ctg gat gaa ggc atc gat      2001
Thr Ser Gln Val Gln Leu Ser Glu Leu Leu Leu Asp Glu Gly Ile Asp
35                  40                  45                  50

atc acc cag gcc acc ttg tcc cgg gat ctc gat gaa ctc ggt gca cgc      2049
Ile Thr Gln Ala Thr Leu Ser Arg Asp Leu Asp Glu Leu Gly Ala Arg
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aag gtt cgc ccc gat ggg gga cgc gcc tac tac gcg gtc ggc cca gta      2097
Lys Val Arg Pro Asp Gly Gly Arg Ala Tyr Tyr Ala Val Gly Pro Val
            70                  75                  80

gat agc atc gcc cgc gaa gat ctc cgg ggt ccg tcg gag aag ctg cgc      2145
Asp Ser Ile Ala Arg Glu Asp Leu Arg Gly Pro Ser Glu Lys Leu Arg
        85                  90                  95

cgc atg ctt gat gaa ctg ctg gtt tct aca gat cat tcc ggc aac atc      2193
Arg Met Leu Asp Glu Leu Leu Val Ser Thr Asp His Ser Gly Asn Ile
    100                 105                 110

gcg atg ctg cgc acc ccg ccg gga gct gcc cag tac ctg gca agt ttc      2241
Ala Met Leu Arg Thr Pro Pro Gly Ala Ala Gln Tyr Leu Ala Ser Phe
115                 120                 125                 130

atc gat agg gtg ggg ctg aaa gaa gtc gtt ggc acc atc gct ggc gat      2289
Ile Asp Arg Val Gly Leu Lys Glu Val Val Gly Thr Ile Ala Gly Asp
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gac acc gtt ttt gtt ctc gcc cgt gat ccg ctc aca ggt aaa gaa cta      2337
Asp Thr Val Phe Val Leu Ala Arg Asp Pro Leu Thr Gly Lys Glu Leu
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ggt gaa tta ctc agc ggg cgc acc act taaagcgccc ctagttcaag           2384
Gly Glu Leu Leu Ser Gly Arg Thr Thr
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Lys Val Thr Ser Gln Val Gln Leu Ser Glu Leu Leu Leu Asp Glu Gly
35 40 45

Ile Asp Ile Thr Gln Ala Thr Leu Ser Arg Asp Leu Asp Glu Leu Gly
50 55 60

Ala Arg Lys Val Arg Pro Asp Gly Gly Arg Ala Tyr Tyr Ala Val Gly
65 70 75 80

Pro Val Asp Ser Ile Ala Arg Glu Asp Leu Arg Gly Pro Ser Glu Lys
85 90 95

Leu Arg Arg Met Leu Asp Glu Leu Leu Val Ser Thr Asp His Ser Gly
100 105 110

Asn Ile Ala Met Leu Arg Thr Pro Pro Gly Ala Ala Gln Tyr Leu Ala

115 120 125

Ser Phe Ile Asp Arg Val Gly Leu Lys Glu Val Val Gly Thr Ile Ala
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